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Fig.1A.

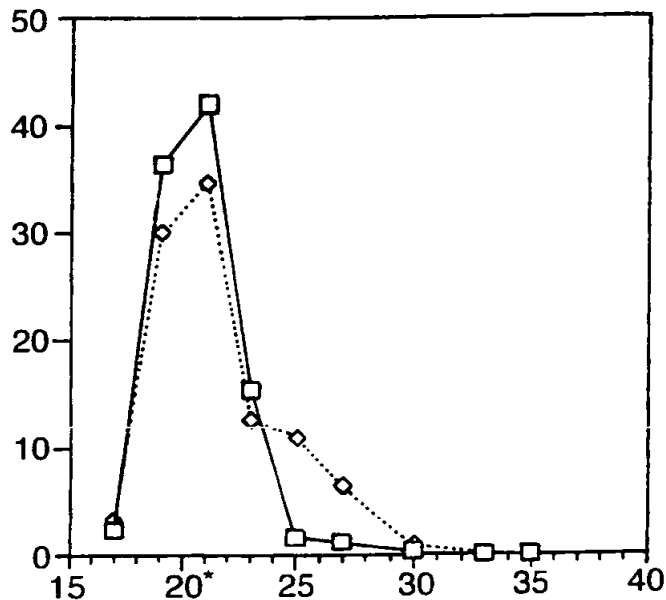
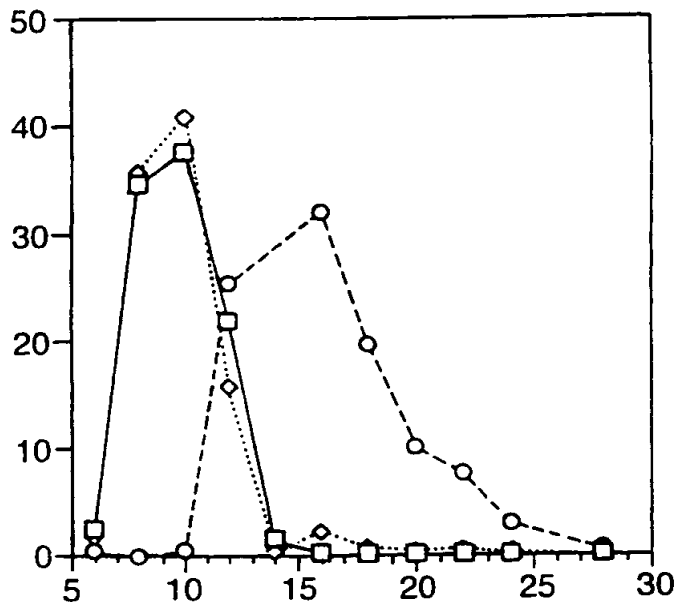


Fig.1B.



662740" 505T4E60

1 HISALOSIPERONFAPSDFKWLCEZELPVKINEVQINGTACTCKSRSEFYELISHVZHRKTVGNNIYPALPTDERRININIDYVETATISGAYKDPKMSATSONKDNKORVQ
 1HNCIDIPOLZREHMAICITETMBARKLIIELENGERDOKDALKUNYRTPTC
 121 KCONLSLLVER..IARRBAEPSSKAIETIDNTHNYDELSQDRFSORPSSVMSPTFHCVENHSPVELRTFFDIVLKHVRVIGQOSUKLLNCMPORODYLSISDSVPTAKYDOK
 51 THDAGOTABINAYPULPCLQKC.SLSLQOQWDLGDIASNNKRR...RQDIA..LSLGLITQSSALCQQLIRHIIOLKLCQVSQQITPSVTHDARLNNNTTDLKXKQCHQCHOPPS
 239 VRKDDOLGIVKVCPTAPQONKKVNLSTKICRTL.DODLVEBEMDORIKQVYMYGSIINFERMOIDITLYLQBSLQSTISQHLR..PTDSVKEGVUGEDIVTPDAKRVILPTC
 166 VCG..SLSLTLPSASKCHMAIADI..SHKEDMXJOSPYIEICLDERMOMH...KDDVYKYSNOMYNDOPGIPTEQSLTPFINNAKADIQICIDOCEDIMAYNPNTOTTPKQK
 356 LVKOSAKZALSPNSINNVPHPLEHVVZDLYTQCTTPTPLPHQKQYDNGHLSBLKNIIVFVRSRRCYGVZSIKKSEVTLISLOSUVLYYNGSIRVAANNHNIIRVREXVRESFOR
 280 TKPOIKRH.....VEDSLOTCCVZQYCHVGNKKCHETKKEIETISHTIPCRIDVYQKTQANTKNEVIDANEYDKRESQINVLSQSLIKPDKGORGOLKREXVSOIAD
 476 NODLIVIG..ROSGKXDSPTLOLVLDSEYTHQDSSZVVDHSSOZXHIGNSRRVKKILSPGZIANQISQEPKZIDQTRONKRRSEVUTPAQIDSPGSKIGANWQDSEKYLE
 394 EMDILIVQCYHUKGRCCHSH7CAYAG..YVQNGPFDTCVLAOSENIRVKNILSNKHVVYKBAWDEGFKTSFVPMQWTHIHICPSTKHTARRYYOCYHDSYITDOLNQKKEVTSOIKN
 594 13SRRLDNTETNHOKTAPNCEIYGCYCKRIRYDKWZDGYETINDYZSRTVKS.....NPSYQASRLOLIRKAKRVLSDEPHQKQOL.PISHTIAGLLIYVTSYVTSAD
 492 13AAEIVPD...HUKGCEIATPPRIEX18DQXWHECH14DDDEQLROKASCKLASKHLVIGDDEPOCKKXKAAPKMKVIGCIEHLKAPDLTNVKNISHTIPEDVPCVMDGOTDAQ.
 702 TQIRITRAEOKETIVHCCGLIYNVTLKXNSIGDVRLECKTTTCAL..DROYILNEMWQDGIAYGRLILIEYNYCTNVSQKXAAVAKRVYOCGLGDSFENDIETKSSLYKSQLS
 607PKPOCENR1A3PCC.....YVQNGPFDTCVLAOSENIRVKNILSNKHVVYKBAWDEGFKTSFVPMQWTHIHICPSTKHTARRYYOCYHDSYITDOLNQKKEVTSOIKN
 821 LPPKCELQIGQVRRP.....13P13SNRIATVER.....RKISTEDDIIEKHLIPGOKITDOOSLCNLIIIPYTDPILRKDCMNGWTHKXKQKASDIPKXIRVYVAPSDPHS
 717 SNEQITPEKALADIADLEIRYSWDC13SMDRRHIVLDSYAVINDLQKXKCTH1A14ALELPHOKN.....VVSCLAEQVSHVIOGCHSRVADPKAPRNPKNKPKILKREHVTDS
 928 1NENQVPPEDPPVNYT.
 831 1DKCELOE3NOYLI.

Figure 2

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Fig.3A.

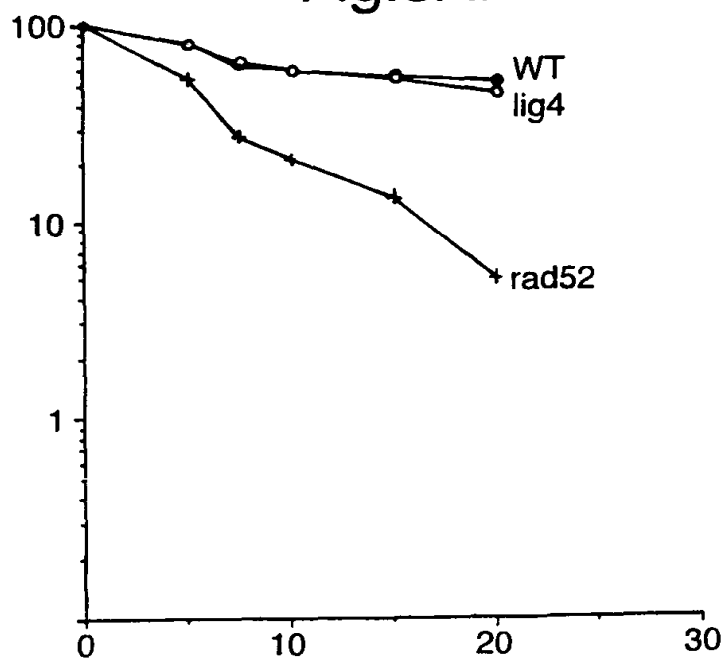
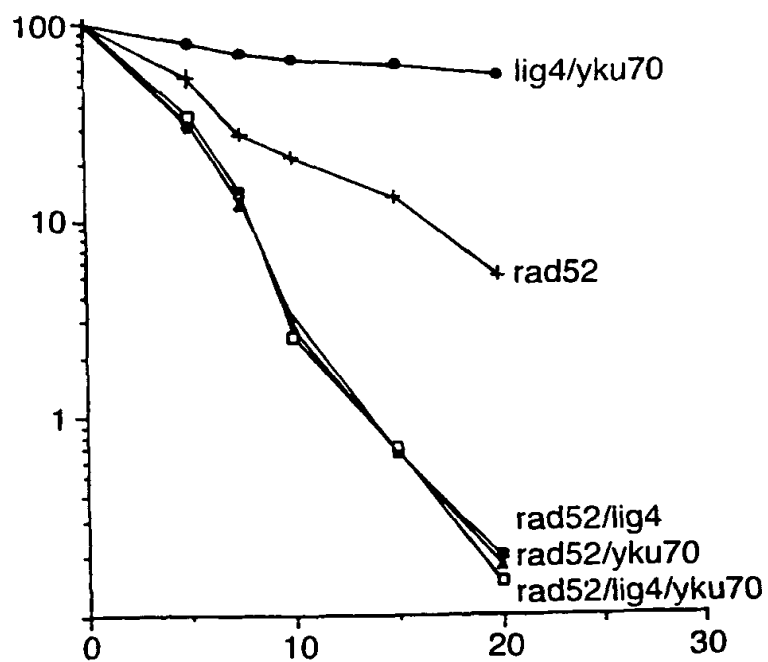
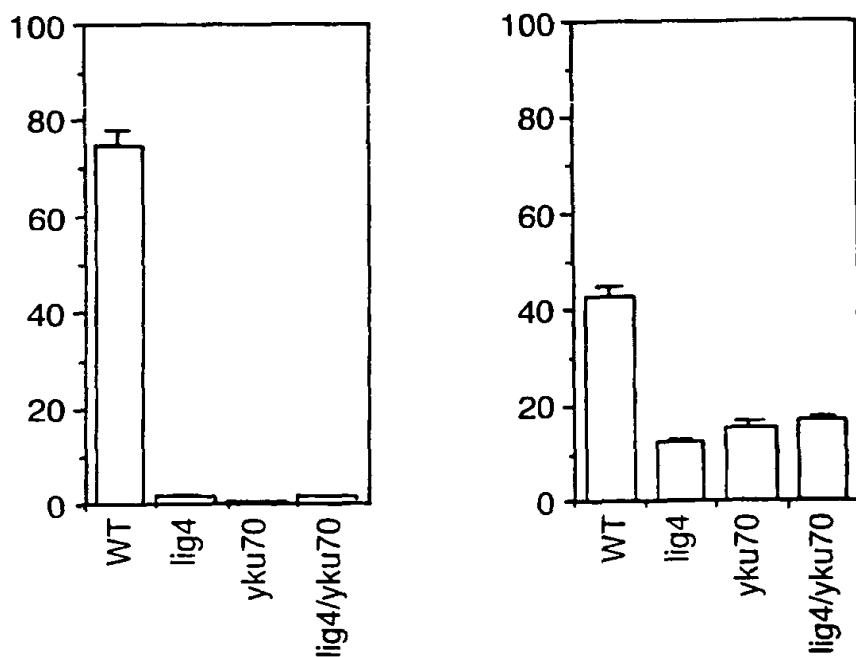


Fig.3B.



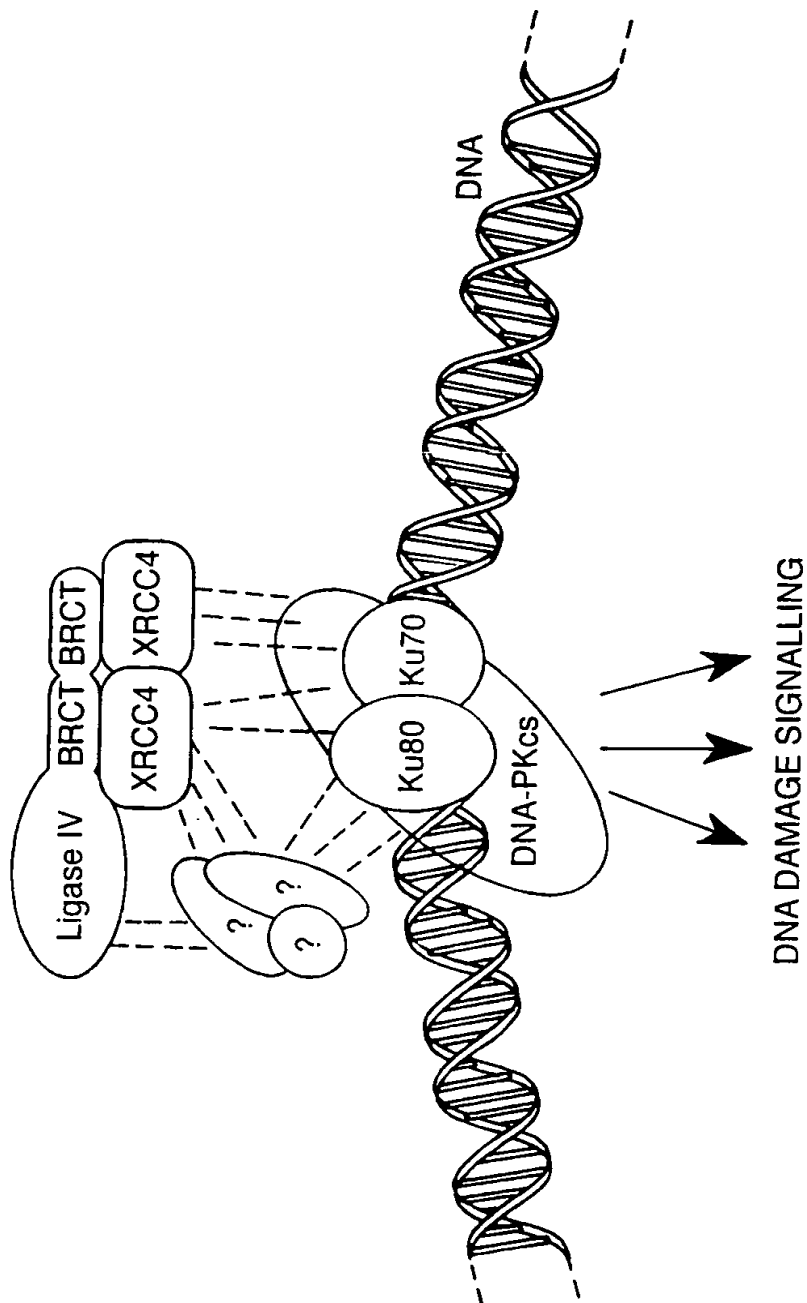
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Fig.4.



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Fig.5.



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Figure 6

GTGATTAAATAGGCTGAAATCAGTGTCTAGTAACTACGTACGTTGTACATGTAAACATTGT
 1 -----+-----+-----+-----+-----+ 60
 CACTAATTTATCCGACTTTAGTCACAAATCATTGATGCATGCAACATGTACATTGTAAACA
 V I K * A E I S V * * L R T L Y M * H C -
 GATATAAATCGTAAGATTGCGCGAGTATAGATCAATAATATCGGTTTCATCACTTACGTT
 61 -----+-----+-----+-----+-----+ 120
 CTATATTTAGCATTCTAAGCGGCTCATATCTAGTTATTATAGCCAAAGTAGTGAATGCAA
 D I N R K I R R V * I N N I G F I T Y V -
 GTTGTGCACTACTAGAGTTAAGATCGTTTCGATCCCTTATTTCTTCTTTTTCCTTT
 121 -----+-----+-----+-----+-----+ 180
 CAAACACGTCATGATCTCAATTCTAGCAAAAGCTAGGGAATAAAGAAGAAAAAAGGAAA
 V C A V L E L R S F S I P Y F L L F S F -
 TTTTGTATTTTCTCTTTTACCTTTTGTCAACCATATTAATCTTTTAAACAAATCTAA
 181 -----+-----+-----+-----+-----+ 240
 AAAAAAATAAAGAGAAAAATGGAACAGTGGTATAATTTAGAAATTTGTTTAGATT
 F L L F F S F Y L L S P Y * I F K Q I * -
 CTATGAAAAAATCCTTTAAACATATGTTAATATGTGGAAATAAATACTAAATAAATAAT
 241 -----+-----+-----+-----+-----+ 300
 GATACTTTTTTAGGAAATTTGTATACAATTATACACCTTTTATTTATGATTTTATTTTA
 L * K N P L N I C * Y V E N K Y * N K N -
 CTAGAAGTGAAGGAAATAGTAACGGATTATTTAGGTATGATATCAGCACTAGATTCTATA
 301 -----+-----+-----+-----+-----+ 360
 GATCTTGACTTCCTTTATCATGCTTAATAAATCCATACTATAGTCGTGATCTAAGATAT
 L E L K E I V T D Y L G M I S A L D S I -
 CCGAGCCCCAAAACCTTTGCGCCTAGTCCAGATTTCAAATGGCTTTGTGAAGAGCTATTT
 361 -----+-----+-----+-----+-----+ 420
 GGGCTCGGGTTTTTGAAACGCGGATCAGGTCTAAAGTTTACCGAAACACTTCTCGATAAA
 P E P Q N F A P S P D F K W L C E E L F -
 GTGAAGATACATGAAGTTCAAAATTAATGGAACGGCGGCACTGGCAAATCAAGGTCTTTC
 421 -----+-----+-----+-----+-----+ 480
 CACTTCTATGTACTTCAAGTTTAATTACCTTGCGGCGGTGACCGTTTAGTTCCAGAAAG
 V K I H E V Q I N G T A G T G K S R S F -
 AAGTACTATGAAATAATATCGAATTTTCGTGAAATGTGGAGAAAAACCGTGGGAAATAAT
 481 -----+-----+-----+-----+-----+ 540
 TTCATGATACTTTATTATAGCTTAAAGCAGCTTTACACCTCTTTTGGCACCCCTTTATTA
 K Y Y E I I S N F V E M W R K T V G N N -
 ATATATCCTGCACTGGTTCTTGCTCTTCCCTACCGGATAGACGAATCTATAATATTAAG
 541 -----+-----+-----+-----+-----+ 600
 TATATAGGACGTGACCAAGAACGAGAAGGGATGGCGCTATCTGCTTAGATATTATAATTC
 I Y P A L V L A L P Y R D R R I Y N I K -

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Figure 6 (Continued)

601 GATTATGTATTAAATAAGAACTATATGCTCTTACTTGAAGTTGCCAAAAAATTCTGCAACA
-----+-----+-----+-----+-----+ 660
CTAATACATAATTATCTTGATATACGAGAATGAACCTCAACGGTTTTTAAAGACGTGT
D Y V L I R T I C S Y L K L P K N S A T -
661 GAGCAGCGGTTAAAGATTGGAACAGCGTGTCCGTAAGGTGGGAATCTTTCTTCTT
-----+-----+-----+-----+-----+ 720
CTCGTCGCCAATTTTCTAACCTTTGTGCGACAGCCATTTCACCCTTAGAAAGAAGAGAA
E Q R L K D W K Q R V G K G G N L S S L -
721 CTGTGTGAAGAAATTGCTAAAAGAAGGGCTGAACCTAGCTCAAAAGCGATTACAATTGAT
-----+-----+-----+-----+-----+ 780
GAACACCTTCTTTAAGGATTTTCTTCCCGACTTGGATCGAGTTTTCGCTAATGTTAACTA
L V E E I A K R R A E P S S K A I T I D -
781 AACGTCAATCACTATCTGGATAGTTTGAAGTGGAGACAGGTTCCGTTCCGGACGAGGATTT
-----+-----+-----+-----+-----+ 840
TTGCAGTTAGTGATAGACCTATCAAACCTCACCTCTGTCCAAGCGAAGGCTGCTCTCTAAA
N V N H Y L D S L S G D R F A S G R G F -
841 AAGAGTCTTGTCAAGTCCAAACCTTTCTGCACTGTGTGGAGAATATGAGTTTCTGTCGAA
-----+-----+-----+-----+-----+ 900
TTCTCAGAACAGTTTCAAGTTTGGAAAGGACGTGACACACCTCTTATACTCAAAGCAGCTT
K S L V K S K P F L H C V E N M S F V E -
901 TTAAATACTTCTTTGATATCGTCTTAAAAATAGAGTAATAGGAGGTCAAGAGCACAAA
-----+-----+-----+-----+-----+ 960
AATTTTATGAAGAACTATAGCAAGATTTTATCTCATTATCCTCCAGTTCTCGTGT
L K Y F F D I V L K N R V I G G Q E H K -
961 TTGCTAAACTGCTGGCATCCTGATGCTCAGGATTATCTTAGCGTGATATCTGATTTAAAG
-----+-----+-----+-----+-----+ 1020
AACGATTTGACGACCGTAGGACTACGAGTCCTAATAGAATCGCACTATAGACTAAATTC
L L N C W H P D A Q D Y L S V I S D L K -
1021 GTGGTAACTTCAAACTTTATGATCCAAAAGTTCTGCTAAAAGGATGATGATTTGAGTATA
-----+-----+-----+-----+-----+ 1080
CACCATTGAAGTTTGAATACTAGGTTTTCAAGCAGATTTCCTACTACTAAACTCATAT
V V T S K L Y D P K V R L K D D D L S I -
1081 AAAGTTGGCTTTGCATTCGCCCCCAATTAGCCAAAAAAGTGAATCTTTCTTATGAGAAA
-----+-----+-----+-----+-----+ 1140
TTTCAACCGAAACGTAAGCGGGGGTTAATCGGTTTTTCACTTAGAAAGAATACTCTTT
K V G F A F A P Q L A K K V N L S Y E K -
1141 ATATGCCGTACACTACATGATGATTTTTTGGTAGAAGAAAAAATGGATGGAGAACGAATT
-----+-----+-----+-----+-----+ 1200
TATACGGCATGTGATGTACTACTAAAAAACCATCTTCTTTTTTACCTACCTCTTGCTTAA
I C R T L H D D F L V E E K M D G E R I -
1201 CAAGTTCATTATATGAATTATGGTGAATCCATAAAATTTTTTAGTAGACGGGCATCGAC
-----+-----+-----+-----+-----+ 1260
GTTCAAGTAATATACTTAATACCACTTAGGTATTTTTAAAAAATCATCTGCCCCGTAGCTG
Q V H Y M N Y G E S I K F F S R R G I D -

0341505 01299 662120 5054450

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Figure 6 (Continued)

1261 TATACCTATTGTGACGAGCGAGCTTATCATCAGGAACATATCTCAACATTTGAGGTTT
 -----+-----+-----+-----+-----+-----+-----+ 1320
 ATATGGATAAACATGCCTCGCTCGAATAGTAGTCCTTGATATAGAGTTGTAAACTCCAAA

 Y T Y L Y G A S L S S G T I S Q H L R F -

 1321 ACAGATAGTGTTAAAGAATGTGTTTTAGATGGAGAAATGGTGACGTTTGATGCAAAAAGA
 -----+-----+-----+-----+-----+-----+-----+ 1380
 TGTCTATCACAATTTCTTACACAAAATCTACCTCTTTACCACTGCAAACTACGTTTTTCT

 T D S V K E C V L D G E M V T F D A K R -

 1381 CGGGTGATTCTTCCATTCCGGTCTTGTAAAGGAAGTGCAAAGGAAGCGCTATCTTTTAAT
 -----+-----+-----+-----+-----+-----+-----+ 1440
 GCCCACTAAGAAGGTAAGCCAGAACAAATTTCTTCACGTTTCTTCGCGATAGAAAATTA

 R V I L P F G L V K G S A K E A L S F N -

 1441 AGTATAAATAATGTTGACTTTCACCCCTTATATATGGTGTTTGATCTGTTATACCTGAAT
 -----+-----+-----+-----+-----+-----+-----+ 1500
 TCATATTTTATTACAACGTGAAGTGCGGAATATATACCACAACTAGACAATATGGACTTA

 S I N N V D F H P L Y M V F D L L Y L N -

 1501 GGGACTTCGTTGACACCATTACCCCTTCATCAAAGGAAGCAATATCTGAACAGCATTTTA
 -----+-----+-----+-----+-----+-----+-----+ 1560
 CCCTGAAGCAACTGTGTAATGGGGAAGTAGTTTCCTTCGTTATAGACTTGTCTGTAAT

 G T S L T P L P L H Q R K Q Y L N S I L -

 1561 AGTCCCTTGAAAAATATGTAGAAATAGTACGATCTTCTAGATGTTATGGTGTTGAGTCA
 -----+-----+-----+-----+-----+-----+-----+ 1620
 TCAGGGAACCTTTTATAACATCTTTATCATGCTAGAAGATCTACAATACCACACCTCAGT

 S P L K N I V E I V R S S R C Y G V E S -

 1621 ATCAAAAAGTCTTTAGAAGTTGCAATCTCACTGGGTTGAGAAGGAGTTGTTTTGAAATAT
 -----+-----+-----+-----+-----+-----+-----+ 1680
 TAGTTTTTCAGAAATCTTCAACGTTAGAGTGACCAAGTCTTCCTCAACAAAACCTTTATA

 I K K S L E V A I S L G S E G V V L K Y -

 1681 TATAATTCAGTTATAATGTGCGCAGTCGAAACAACAACTGGATCAAGGTAAAACCTGAA
 -----+-----+-----+-----+-----+-----+-----+ 1740
 ATATTAAGTTCAATATTACAGCGGTCAGCTTTGTTGTTGACCTAGTTCCATTTTGGACTT

 Y N S S Y N V A S R N N N W I K V K P E -

 1741 TATTTGGAGGAATTTGGAGAGAATTTAGACTTAATAGTAATAGGCAGAGATTCTGGGAAA
 -----+-----+-----+-----+-----+-----+-----+ 1800
 ATAAACCTCCTTAAACCTCTCTTAAATCTGAATTATCATTATCCGTCTCTAAGACCCCTT

 Y L E E F G E N L D L I V I G R D S G K -

 1801 AAAGATTCTTTTATGCTAGGGTTACTTGTGCTAGATGAAGAAGAGTATAAAAAGCACCAA
 -----+-----+-----+-----+-----+-----+-----+ 1860
 TTTCTAAGAAAATACGATCCCAATGAACACGATCTACTTCTTCTCATATTTTTCGTGGTT

 K D S F M L G L L V L D E E E Y K K H Q -

 1861 GGAGACTCCTCTGAAATTGTAGACCACTCAAGCCAAGAAAAACACATACAAAATTCAAGA
 -----+-----+-----+-----+-----+-----+-----+ 1920
 CCTCTGAGGAGACTTTAACATCTGGTGAGTTCCGTTCTTTTGTGTATGTTTTAAGTTCT

 G D S S E I V D H S S Q E K H I Q N S R -

 AGAAGGGTGAAAAAATACTTTCATTCTGTTCTATCGCAAACGGTATATCTCAAGAGAA

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Figure 6 (Continued)

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1921 -----+-----+-----+-----+-----+-----+ 1980
      TCTTCCCACTTTTTTTATGAAAGTAAGACAAGATAGCGTTTGCCATATAGAGTTCTTCTT
      R R V K K I L S F C S I A N G I S Q E E -
      TTCAAAGAAATCGACCGCAAAACGAGAGGACATTGGAAAAGAACCTCCGAAGTTGCTCCC
1981 -----+-----+-----+-----+-----+ 2040
      AAGTTTCTTTAGCTGGCGTTTTGCTCTCCGTAAACCTTTTCTTGGAGGCTTCAACGAGGG
      F K E I D R K T R G H W K R T S E V A P -
      CCTGCTTCAATTTTAGAATTTGGCTCAAAAATACCTGCCGAATGGATTGACCCCAGTGAA
2041 -----+-----+-----+-----+-----+ 2100
      GGACGAAGTTAAATCTTAAACGAGTTTTTATGGACGGCTTACCTAACTGGGGTCACTT
      P A S I L E F G S K I P A E W I D P S E -
      TCAATTGTTCTAGAAATAAAATCACGGTCTTTGGATAACACAGAAACGAATATGCAGAAG
2101 -----+-----+-----+-----+-----+ 2160
      AGTTAACAAGATCTTTATTTTAGTGCCAGAAACCTATTGTGCTTTTGCTTATACGCTCTC
      S I V L E I K S R S L D N T E T N M Q K -
      TACGCTACCAATTGTACTTTGTACGGTGGCTATTGTAAAAGAATACGGTACGATAAAGAA
2161 -----+-----+-----+-----+-----+ 2220
      ATGCGATGGTTAACATGAAACATGCCACCGATAACATTTTCTTATGCCATGCTATTCTT
      Y A T N C T L Y G G Y C K R I R Y D K E -
      TGGACAGATTGTTACACACTTAACGACTTATACGAAAGTAGGACGGTTAAATCTAACCCC
2221 -----+-----+-----+-----+-----+ 2280
      ACCTGTCTAACAATGTGTGAATTGCTGAATATGCTTTCATCCTGCCAATTTAGATTGGGG
      W T D C Y T L N D L Y E S R T V K S N P -
      AGCTATCAAGCGGAAAGGTCACAGCTTGGATTGATACGGAAAAAGAGAAAGAGAGTACTT
2281 -----+-----+-----+-----+-----+ 2340
      TCGATAGTTGCGCTTTCCAGTGTGGAACCTAACTATGCCTTTTTCTCTTCTCTCATGAA
      S Y Q A E R S Q L G L I R K K R K R V L -
      ATTTACAGACAGCTTTTACCAAAACAGGAAACAACCTGCCAATTTCAAACATCTTTGCCGGA
2341 -----+-----+-----+-----+-----+ 2400
      TAAAGTCTGTGAAAGTGGTTTTGTCTTTGTGACGGTTAAAGTTGTAGAAACGGCCT
      I S D S F H Q N R K Q L P I S N I F A G -
      TTACTTTTTTATGTTCTCTCTGACTATGTACGGAGGACACTGGAATACGGATTACACGG
2401 -----+-----+-----+-----+-----+ 2460
      AATGAAAAAATACAAGAGAGACTGATACAGTGCTCCTGTGACCTTATGCCTAATGTGCC
      L L F Y V L S D Y V T E D T G I R I T R -
      GCAGAACTTGAAAAAACTATTGTGGAACATGGTGGTAAACTGATATATAATGTAATTTTA
2461 -----+-----+-----+-----+-----+ 2520
      CGTCTTGAACCTTTTTTGATAACACCTGTACCACCATTTGACTATATATTACATTAAAT
      A E L E K T I V E H G G K L I Y N V I L -
      AAACGTCATTCAATTGGGGACGTTGCGTTAATCAGCTGTAAACTACCACGGAATGCAAG
2521 -----+-----+-----+-----+-----+ 2580
      TTTGCAGTAAGTTAACCCTGCAAGCCAATTAGTCGACATTTTGATGGTGCCTTACGTT
      K R H S I G D V R L I S C K T T T E C K -
      GCTTTAATAGATGAGGATATGATATATTGCACOCAAATTGGGTACTGATTGTATAGCA
2581 -----+-----+-----+-----+-----+ 2640

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Figure 6 (Continued)

CGAAATTATCTAGCTCCTATACTATATAACGTGGGTTTAACCCATGAGCTAACATATCGT
 A L I D R G Y D I L H P N W V L D C I A -
 TATAAGAGGCTCATCCTGATCGAGCCCAATTATTGCTTTAACGCTCTCAAAAAATGAGA
 2641 -----+-----+-----+-----+-----+ 2700
 ATATTCTCOGAGTAGGACTAGCTCGGGTTAATAACGAAATGTCAGAGAGTTTTTTACTCT
 Y K R L I L I E P N Y C F N V S Q K M R -
 GCCGTCGCTGAAAAAAGGGTAGATTGTTTGGGTGATAGTTTTGAAAATGACATTTTCGGAA
 2701 -----+-----+-----+-----+-----+ 2760
 CGGCAGCGACTTTTTTCCCATCTAACAAACCCACTATCAAAACTTTTACTGTAAAGCCTT
 A V A E K R V D C L G D S F E N D I S E -
 ACCAAACTGTCATCATTTGTATAAATCACAACCTAAGTCTACCACCGATGGGGGAACCTCGAG
 2761 -----+-----+-----+-----+-----+ 2820
 TGGTTTGACAGTAGTAACATATTTAGTGTGATTGAGATGGTGGCTACCCCTTGAGCTC
 T K L S S L Y K S Q L S L P P M G E L E -
 ATAGATTCTGAGGTTTCGGCGGTTTCCATTATTTTTATTCTCCAACAGGATTGCATACGTA
 2821 -----+-----+-----+-----+-----+ 2880
 TATCTAAGACTCCAAGCCGCAAAGGTAATAAAAAATAAGAGGTTGTCTAACGTATGCAT
 I D S E V R R F P L F L F S N R I A Y V -
 CCACGTCGCAAAATTAGCACAGAAGATGACATTATAGAAATGAAAATTAAGTTGTTTGGT
 2881 -----+-----+-----+-----+-----+ 2940
 GGTGCAGCGTTTTAATCGTGTCTTCTACTGTAATATCTTTACTTTTAATTCAACAAACCA
 P R R K I S T E D D I I E M K I K L F G -
 GGAAAAATAACAGATCAACAGTCACTTTGTAACTTAATAATTATACCATATACTGATCCT
 2941 -----+-----+-----+-----+-----+ 3000
 CCTTTTTTATTGTCTAGTTGTGTCAGTGAAACATTGAATTATTAATATGGTATATGACTAGGA
 G K I T D Q Q S L C N L I I I P Y T D P -
 ATTTTGAGGAAAGACTGCATGAATGAGGTACACGAAAAAATAAAAGAACAAATAAAGGCT
 3001 -----+-----+-----+-----+-----+ 3060
 TAAACTCCTTTCTGACGTACTTACTCCATGTGCTTTTTTATTCTTGTGTTATTTTCGA
 I L R K D C M N E V H E K I K E Q I K A -
 TCTGATACTATACCGAAAAATAGCCAGGGTGGTGGCCCTGAATGGGTGGATCATTCTATT
 3061 -----+-----+-----+-----+-----+ 3120
 AGACTATGATATGGCTTTTATCGTCCCGAGCAACGGGGACTTACCCACCTAGTAAGATAA
 S D T I P K I A R V V A P E W V D H S I -
 AATGAAAACTGTCAAGTGCCTGAAGAAGACTTCCCCGTAGTCAACTACTGATGGTGGCTT
 3121 -----+-----+-----+-----+-----+ 3180
 TTACTTTTGACAGTTCAACGACTTCTTCTGAAGGGGCATCAGTTGATGACTACCACGCAA
 N E N C Q V P E E D F P V V N Y * W C V -
 TTGCGGAGGCTTAATTTTTTTGAAGTTTATTTAATACTATCCTACATATGTACATTAAATA
 3181 -----+-----+-----+-----+-----+ 3240
 AACGCCCTCOGAATTAAAAAATTCAAATAAATTATGATAGGATGTATACATGTAATTTAT
 L R R L N F L K F I * Y Y P T Y V H * I -
 CTTCGGTAAACGTTTATCAATAAGAGTGGGAAGATGCGCAATTATATTCAAAAGATTGGCCA
 3241 -----+-----+-----+-----+-----+ 3300
 GAAGGCATTGCAAAATAGTTATTCTCACCTTCTAAGCGTTAATATAAGTTTTCTAACCGGT

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Figure 6 (Continued)

L P * R L S I R V E D A Q L Y S K D W P -
GTCAATTAACTTAAGGAAAAAAT
3301 -----+-----+--- 3323
CAGTTAATTGAATTCCTTTTAA
V N * L K E K -